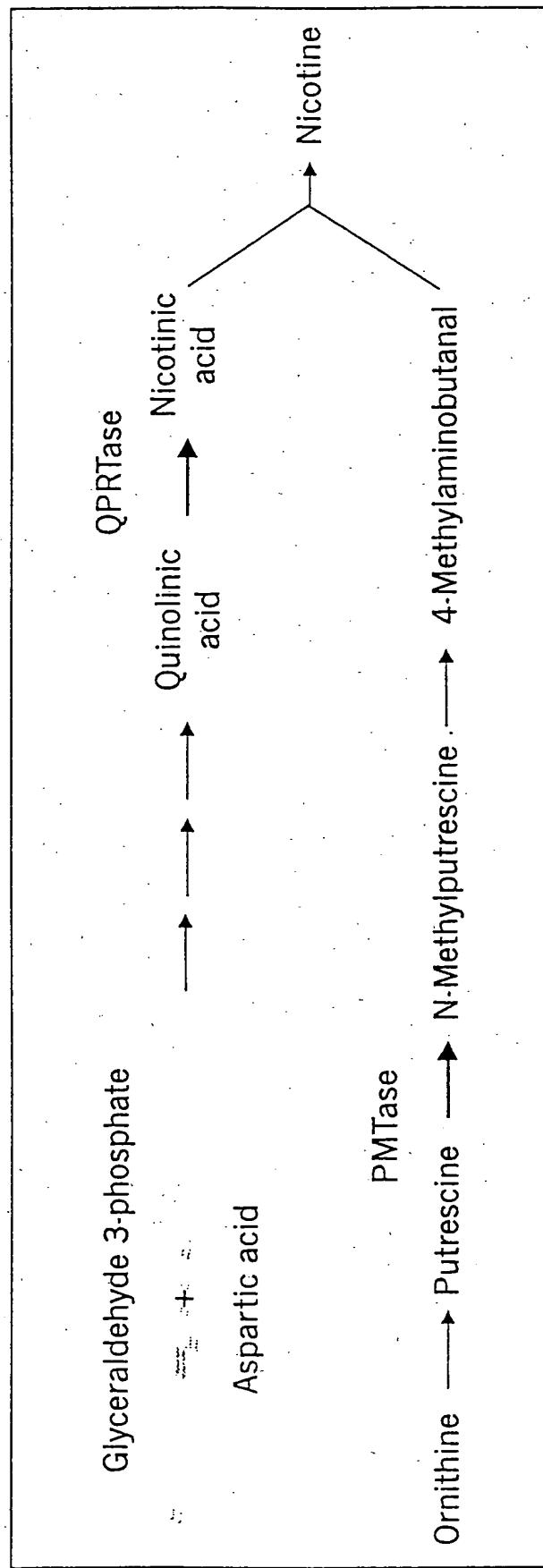


FIG. 1



caaaaactat	tttccacaaa	attcatttca	caacccccc	aaaaaaaaac	CATGTTAGA	60
GCTATTCTT	TCACTGCTAC	AGTCATCCT	TATGCAATT	CAGCTCCAAG	GTTGGTGGTG	120
AAAATGTCAG	CAATAGCCAC	CAAGAATACA	AGAGTGGAGT	CATTAGAGGT	GAAACCACCA	180
GCACACCCAA	CTTATGATT	AAAGGAAGTT	ATGAAACCTG	CACTCTCTGA	AGATGCTGG	240
TTTCTAGCAA	AGGAAGACGG	GATCATAGCA	GGAATTGCAC	TTGCTGAGAT	GATATTGCG	360
GAAGTTGATC	CTTCATTAAA	GGTGGAGTGG	TATGTAATG	ATGGCGATAA	AGTTCATAAA	420
GGCTTGAAAT	TTGGCAAAGT	ACAAGGAAAC	GCTTACAC	TTGTTATAGC	TGAGAGGGTT	480
GTTCTCAATT	TTATGCAAAG	AATGAGTGG	ATAGCTACAC	TAACTAAGGA	AATGGCAGAT	540
GCTGCACACC	CTGCTTACAT	CTGGAGACT	AGGAAAAC	CTCCTGGATT	ACGTTGGTG	600
GATAAATGGG	CGGTATTGAT	CGGTGGGGGG	AAGAATCACA	GAATGGGCTT	ATTTGATATG	660
GTAATGATAA	AAGACAATCA	CATATCTGCT	GCTGGAGGTG	TCGGCAAAGC	TCTAAAATCT	720
GTGGATCAGT	ATTTGGAGCA	AAATAAAACTT	CAAATAGGGG	TTGAGGTTGA	AACCAGGACA	780
ATTGAAGAAG	TACGTGAGGT	TCTAGACTAT	GCATCTCAA	CAAAGACTTC	GTTGACTAGG	840
ATAATGCTGG	ACAATATGGT	TGTTCCATTA	TCTAACGGAG	ATATTGATGT	ATCCATGCTT	900
AAGGAGGCTG	TAGAATTGAT	CAATGGGAGG	TTTGATAACGG	AGGCTTCAGG	AAATGTTACC	960
CTTGAACACAG	TACACAAAGAT	TGGACAAACT	GGTGTACCT	ACATTCTAG	TGGTGCCCTG	1020
ACGCATTCCG	TGAAAGCACT	TGACATTTC	CTGAAGATCG	ATACAGAGCT	CGCCCTTGAA	1080
GTTGGAAGGC	GTACAAAACG	AGCATGAgcg	ccattacttc	tgcatacggtt	ttggagtaaa	1140
agcagctgaa	tagctgaaag	gtgcaaataa	gaatcatttt	actagttgtc	aaacaaaaga	1200
tccttcactg	tgtatcaaa	caaaaagatg	taaattgctg	gaatatctca	gatggctctt	1260
ttccaaacctt	attgcttgag	ttgtaattt	cattatagct	ttgtttcat	gtttcatgga	1320
atttgttaca	atgaaaatac	ttgattata	agtttgggt	atgtaaaatt	ctgtgttact	1380
tcaaataattt	tgagatgtt					1399

FIGURE 2A

MFRAIPFTAT	VHPYAITAPR	LVVKMSAIAT	KNTRVESLEV	KPPAHPTYDL	50
KEVMKLALSE	DAGNLGDVTC	KATIPLDMES	DAHFLAKEDG	IIAGIALAEM	100
IFAEVDP SLK	VEWYVNDGDK	VHKGLKFGKV	QGNAYNIVIA	ERVVLNFMQR	150
MSGIATLTKE	MADAHPAYI	LETRKTAPGL	RLVDKWAVLI	GGGKNHRMGL	200
FDMVMIKDNH	ISAAGGVGKA	LKSVDQYLEQ	NKLQIGVEVE	TRTIEEVREV	250
LDYASQTKTS	LTRIMLDNMV	VPLSNGDIDV	SMLKEAVELI	NGRFDTEASG	300
NVTLETVHKI	GQTGVTYISS	GALTHSVKAL	DISLKIDTEL	ALEVGRRTKR	350
A					

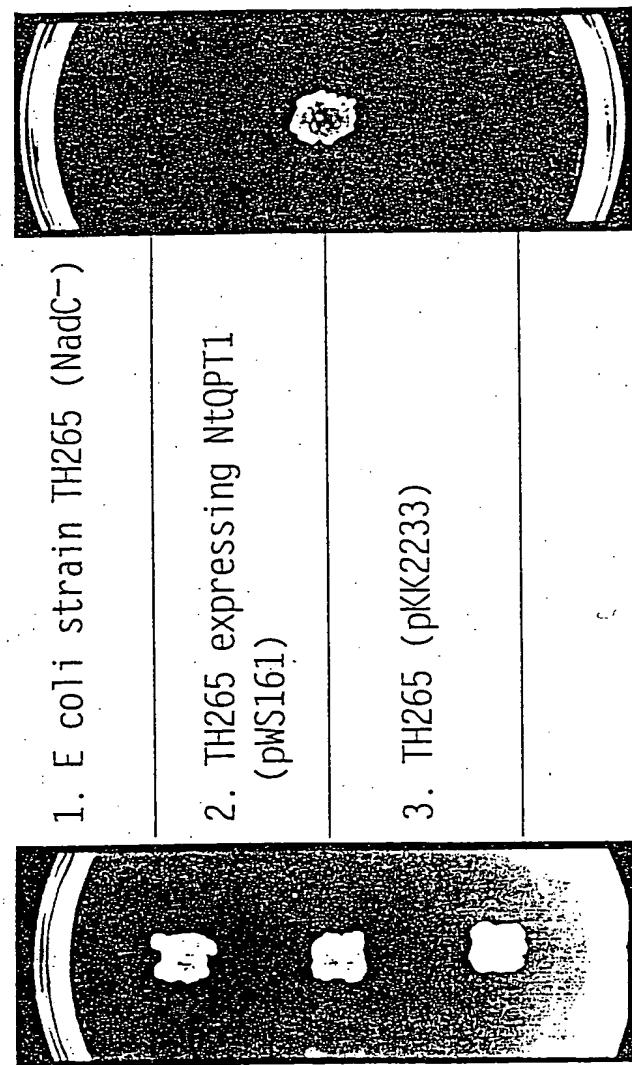
FIGURE 2B

N. tabacum	MFRAIPFTATVHPYAITAPRLVVKMSAIATKNTRVESLEVKKPAHPTYDL
R. rubrum	*-----RPNH-----PVAALS*F---AI
M. leprae	*-----LSDC-----EFDAAR-----
S. typhimurium	*-----PPRR*NPDDR*-----DALL*RINLDI*A---AV
E. coli	*-----PPRR*NPDTR*-----DELL*RINLDI*G---AV
H. sapien	*-----D*EG*ALLLPPVTLAALVDSWLREDC*G-----
S. cerevisiae	*-----PVYE-HLLPVNGAWRQDVTNWLSEDV*S-----
N. tabacum	KEVMKLALSEDAGNLGDTCKATIPLDMESDAHFLAKEDGI IAGIA---
R. rubrum	D*AVRR**A**L*RA**I*ST***AATRAH*RFV*RQP**L**LGCA--
M. leprae	-DTIRR**H**LRYGL*I*TQ**V*AGTVVTGSMVPR*P*VIAGVDALL
S. typhimurium	AQALREDLGGEVDAGN*I*AQL-L*A*TQAH*TVITR*D*VF---CGKR
E. coli	AQALREDLGGTVDANN*I*A*L-L*ENSR*H*TVITR*N*VF---CGKR
H. sapien	-----LNYAALVSGAGP*QAALWAKSP*VL---AGQP
S. cerevisiae	-----FDFGGYVVGSDLKEANLYCKQD*ML---CGVP
N. tabacum	-LAEMIFAEVDPSLKVEWYVNDGDKVHKGLK-----FGKVQGNAYNIVI
R. rubrum	--RSAF-ALLDDTVTFTTPLE**AEIAA*OT-----VAE*A*A*RT*LA
M. leprae	VLD*VF-GVDGYRVLY--R*E**ARLQS*QP-----LLTVQAA*RGLLT
S. typhimurium	WVE*VFIQLAGDDVRLT*H*D***AI*ANQT-----VFELN*PARVLLT
E. coli	WVE*VFIQLAGDDVTI I*H*D***VINANQS-----LFELE*PSRVLLT
H. sapien	FFDAIFTQL---NCQVS*FLPE*S*LVPVAR-----VAEVR*P*HDLLL
S. cerevisiae	FAW*VFNQC---ELQVE*LFKE*SFLEPSKNDSGKIVVAKIT*P*K**LL
N. tabacum	AERVVLFNMORMSGIATLTKEAD--AAH--PAYILETRKTAPGLRLVDK
R. rubrum	***TA***LGHL*****R*RRFG*AI*HT--R*RLTC****T****GLE*
M. leprae	***TM***VCHM*****V*VAWV*AVRGT--K*KIRD****L****ALQ*
S. typhimurium	G**TA***V*TL**VASEVRYYVGLL*GT--QTQL*D****L****TAL*
E. coli	G*PTA***V*TL**VASKVRHYVELLEGT--NTQL*D****L****SAL*
H. sapien	G***A**TLARC****SAAAAAVEAARGAGWTGHVAG****T**F***E*
S. cerevisiae	***TA**ILSRS****TASHKIISLARSTGYKGTIAG****T****RLE*
N. tabacum	WAVLIGGGKNHRMGLFDMVMIKDNHISAAGGVGKALKSVDQYLEQNKLQI
R. rubrum	Y***RC***S***F**D*A*L*****AVA***SA**SRAR-AGVGHMVRI.
M. leprae	Y**RV***V***L**G*TAL*****VA*V*S*VD**RA*R-AAPEL-PC
S. typhimurium	Y***C***A***L**T*AFL*****I*S*S*RQ*VEKAF-W*HPD-APV
E. coli	Y***C***A***L**S*AFL*****I*S*S*RQ*VEKAS-W*HPD-APV
H. sapien	YGL*V**AAS**YD*GGLVML*D**VVPP***EK*VRAARQ---AADFAL
S. cerevisiae	YSM*V**CDT**YD*SS**ML*D***W*T*SITN*V*NARA---VCGFAV

FIGURE 3

N. tabacum	GVEVETRTIEEVREVDYASQTKTSLTRIMLDNMVVPLSNGDIDVSMLE		
R. rubrum	EI****-L*QLA***AVGGAEV----VL***-DAPT---*TR		
M. leprae	E****S--L*QLDAM*A-EEPEL---*L***F--*VWQTQV---AVQ		
S. typhimurium	E****N--LDELDDA*K-*GADI----****F----NTDQ---MR*		
E. coli	E****N--L**LD*A*K-*GADI----****F----ETEQ---MR*		
H. sapien	K****CSSLQ**VQAAE-*GADL----VL***F----KPEELHPTAT		
S. cerevisiae	KI***CLSED*AT*AIE-*GADV----****F----KGDGLK*CAQ		
N. tabacum	AVELI---NGRFDTEASGNVTLETVHKIG-QTGVTYISSGALTHSVKALD		
R. rubrum	**DMV---A**LV*****G*S*D*IAALA-ES**D***V*****TT**		
M. leprae	RRDIR---APTVLL*S**GLS**NAAIYA-G***DYLAV*****RI**		
S. typhimurium	**KRV---**QARL*V*****AE*LREFA-E***DF**VG*****R***		
E. coli	**KRT---**KALL*V*****DK*LREFA-E***DF**V*****Q***		
H. sapien	*LKAQFPSVA---VEA**GIT*DNLPOF-CGPHIDV***M*M**QA*P***		
S. cerevisiae	SLKNKWNGKKHFLLEC**GLN*DNLEYLCD-DIDIY*TSSIHQGTPVI*		
N. tabacum	ISKLIDTELALEVGRRTKRA	<u>% Identity</u>	<u>% Similarity</u>
R. rubrum	*G*D*VVA----PPKAERA	15.9	43.2
M. leprae	*G*DL	18.3	37.3
S. typhimurium	LSMRFC	18.2	34.8
E. coli	LSMRFR	17.9	32.8
H. sapien	F***L---F*K*VAPVP*IH	16.8	31.7
S. cerevisiae	F***LAH	14.6	27.8

FIGURE 3 continued



1. E coli strain TH265 (NadC<sup>-</sup>)

2. TH265 expressing NtQPT1  
(pMS161)

3. TH265 (pKK2233)

ME Minimal Media

ME Minimal Media  
+ Nicotinic Acid

FIG. 4

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FIG. 5

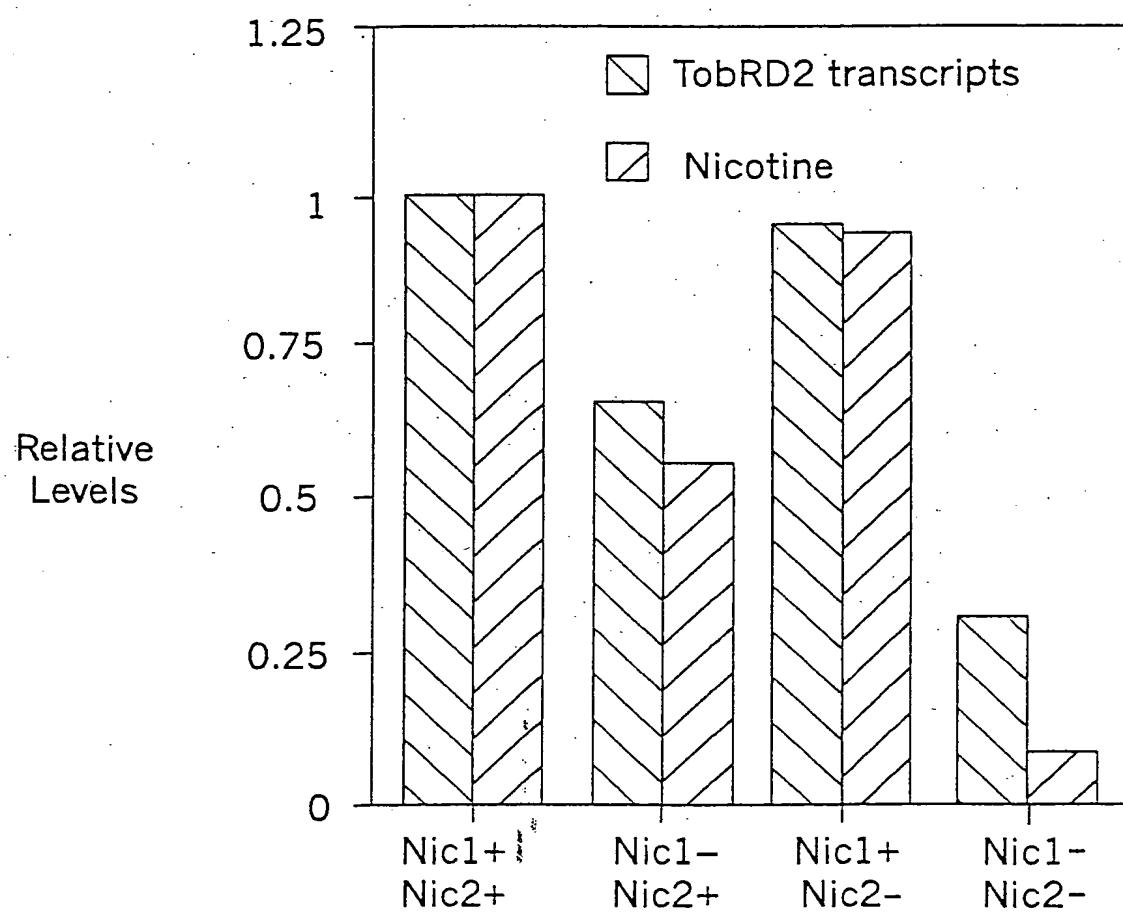


FIG. 6

